

fig. 1A

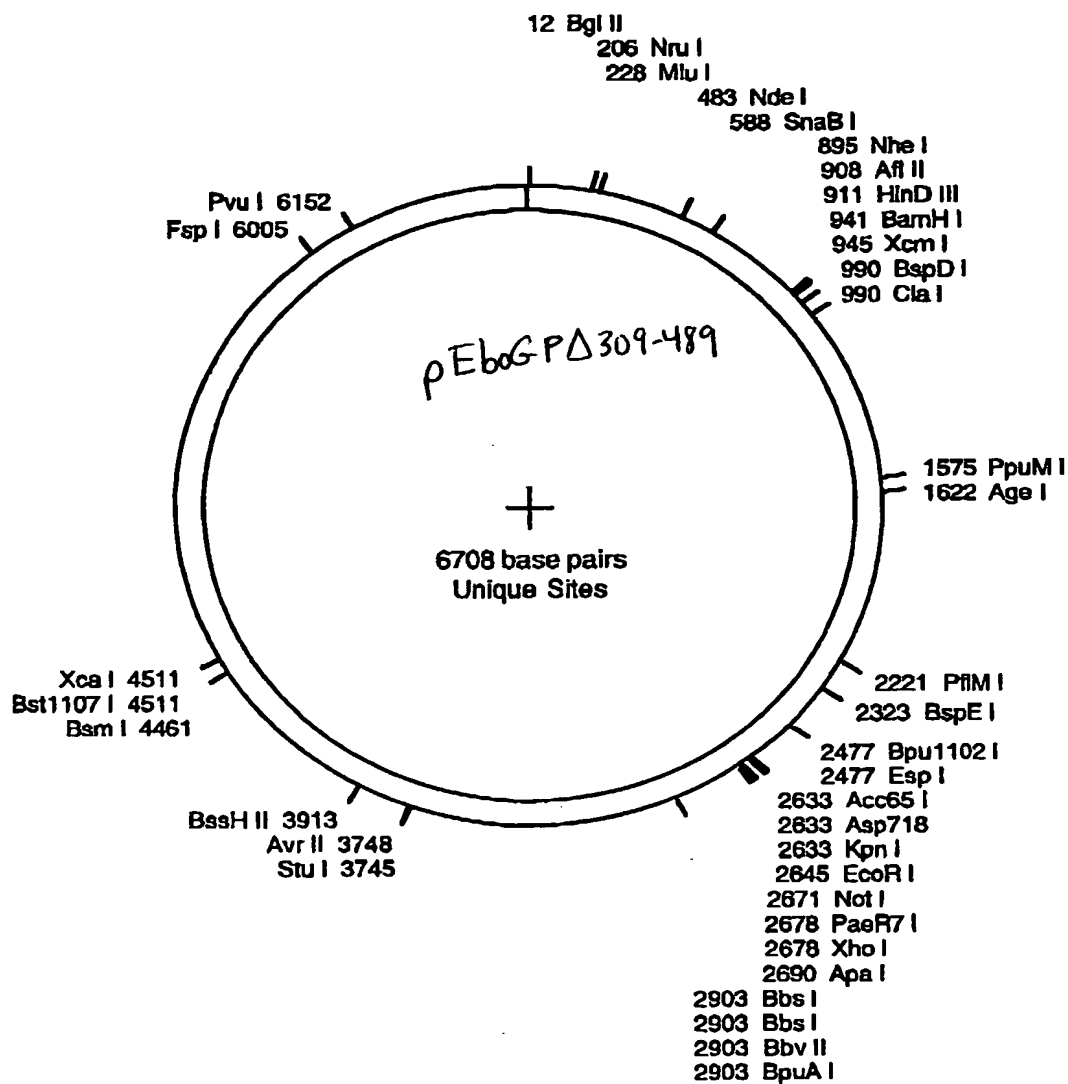
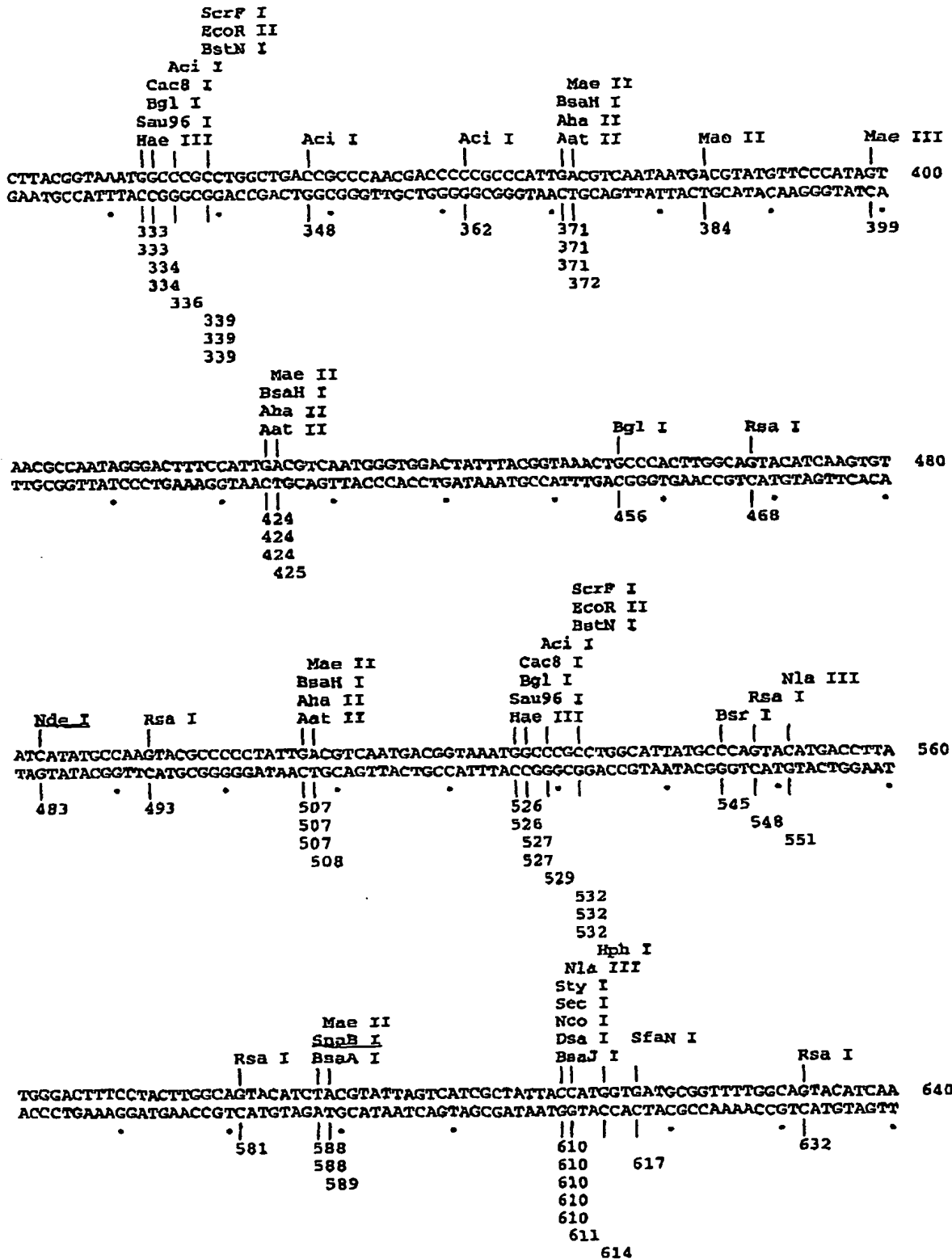
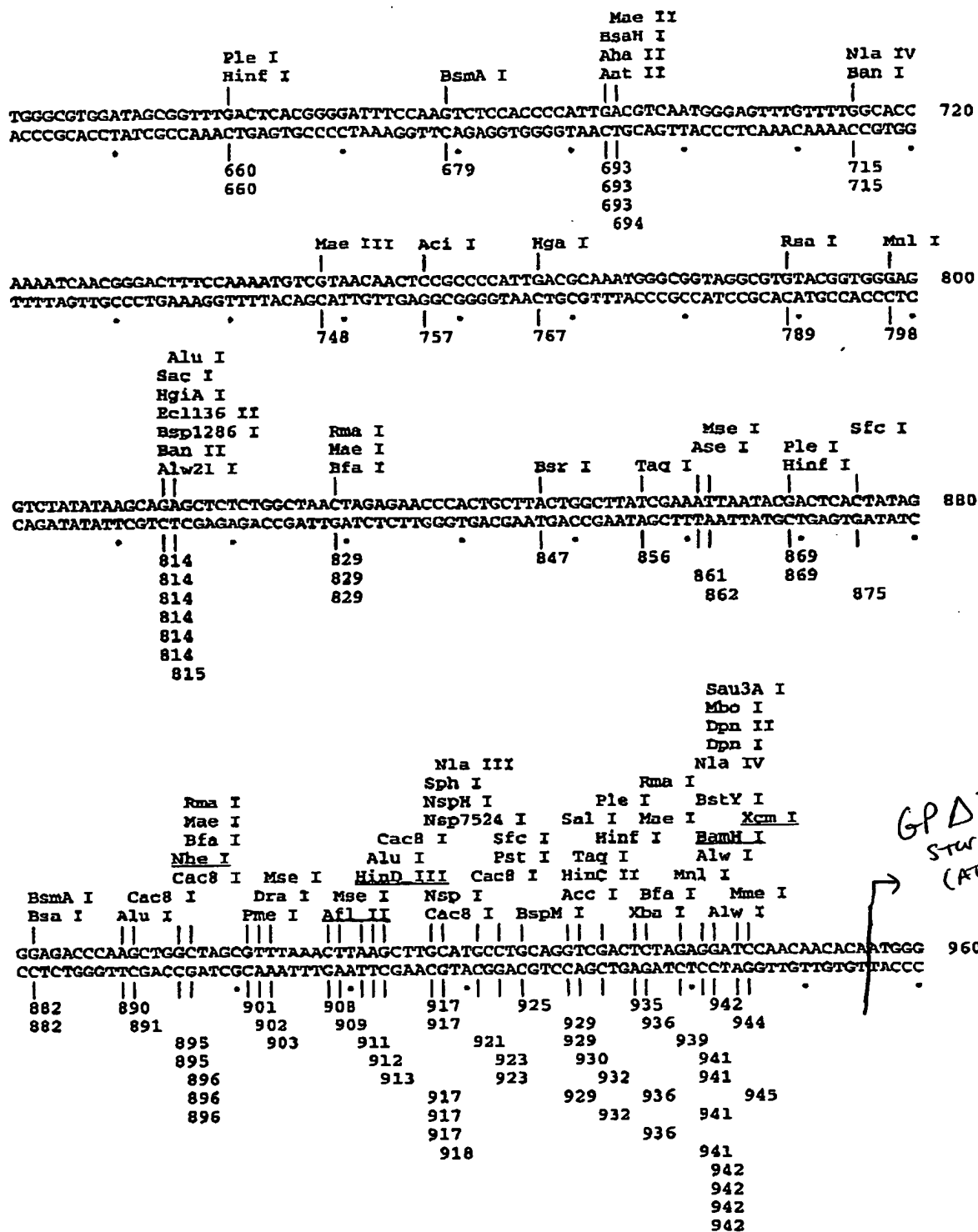


fig. 1B

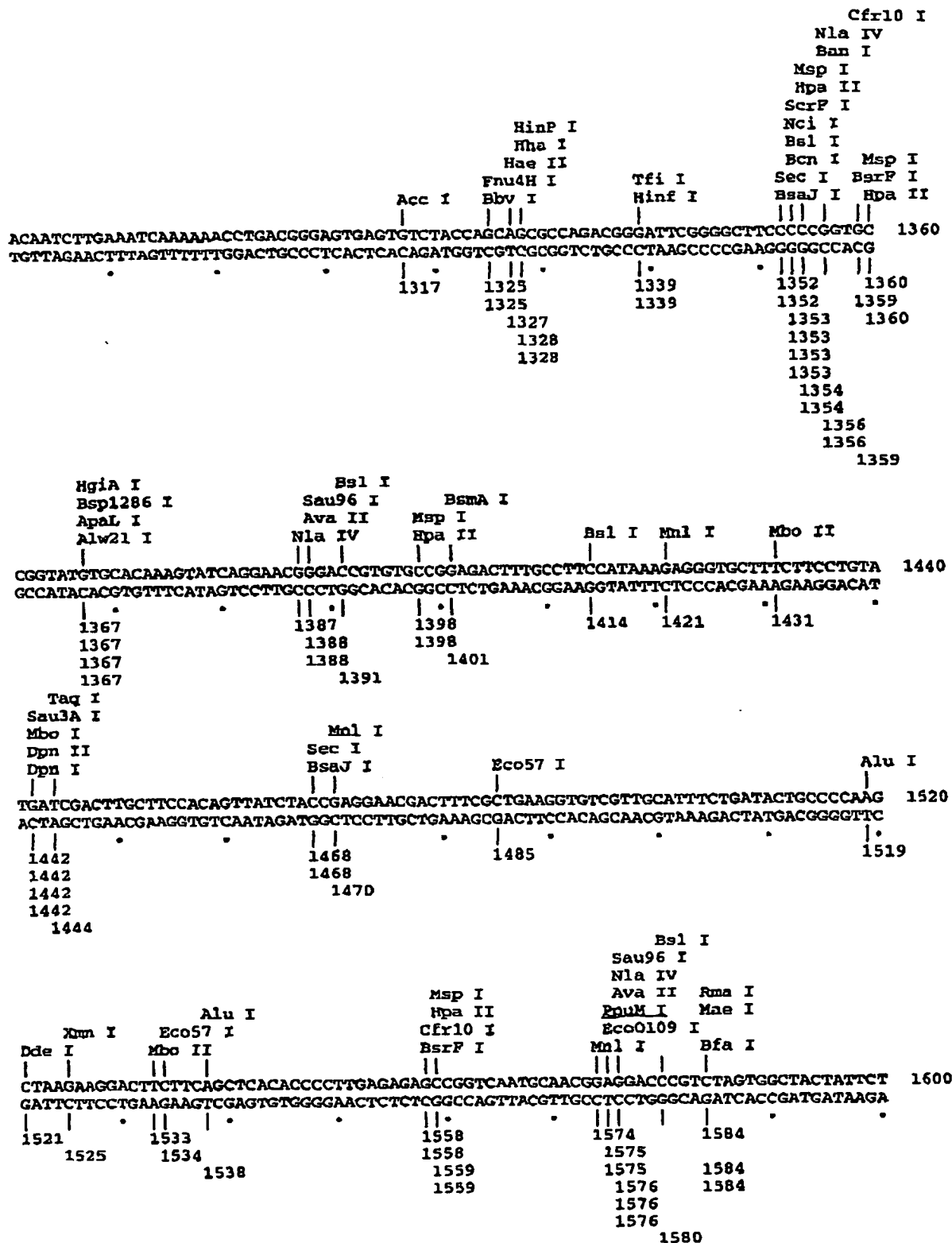
## Positions of Restriction Endonucleases sites (unique sites underlined)

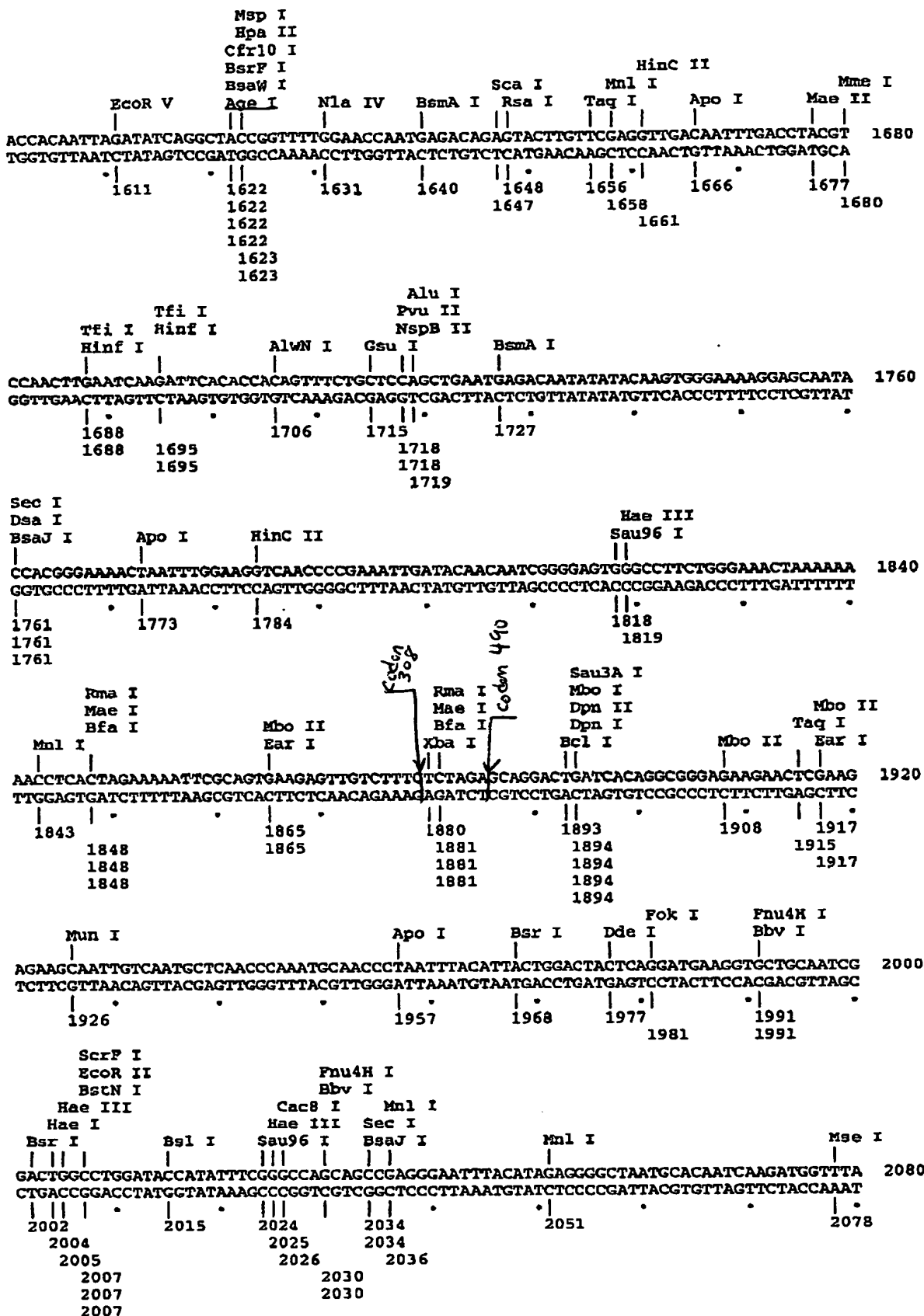
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Mbo I		Dpn II		Dpn I		Dpn II		Hinc II		Dde I		Acc I		Rsa I		Fnu4H I		Bsr I	
Dpn I		Dpn I		Dpn I		Dpn I		Hinc II		Dde I		Acc I		Rsa I		Fnu4H I		Bsr I	
Alw I		Bgl II		Alw I		Alw I		Hinc II		Dde I		Acc I		Rsa I		Fnu4H I		Bsr I	
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4		12		21		32		39		57		69		74		75			
5		12		21		32		43		60		74							
5		13		21		32				61									
5		13		21		33													
5				21		35													
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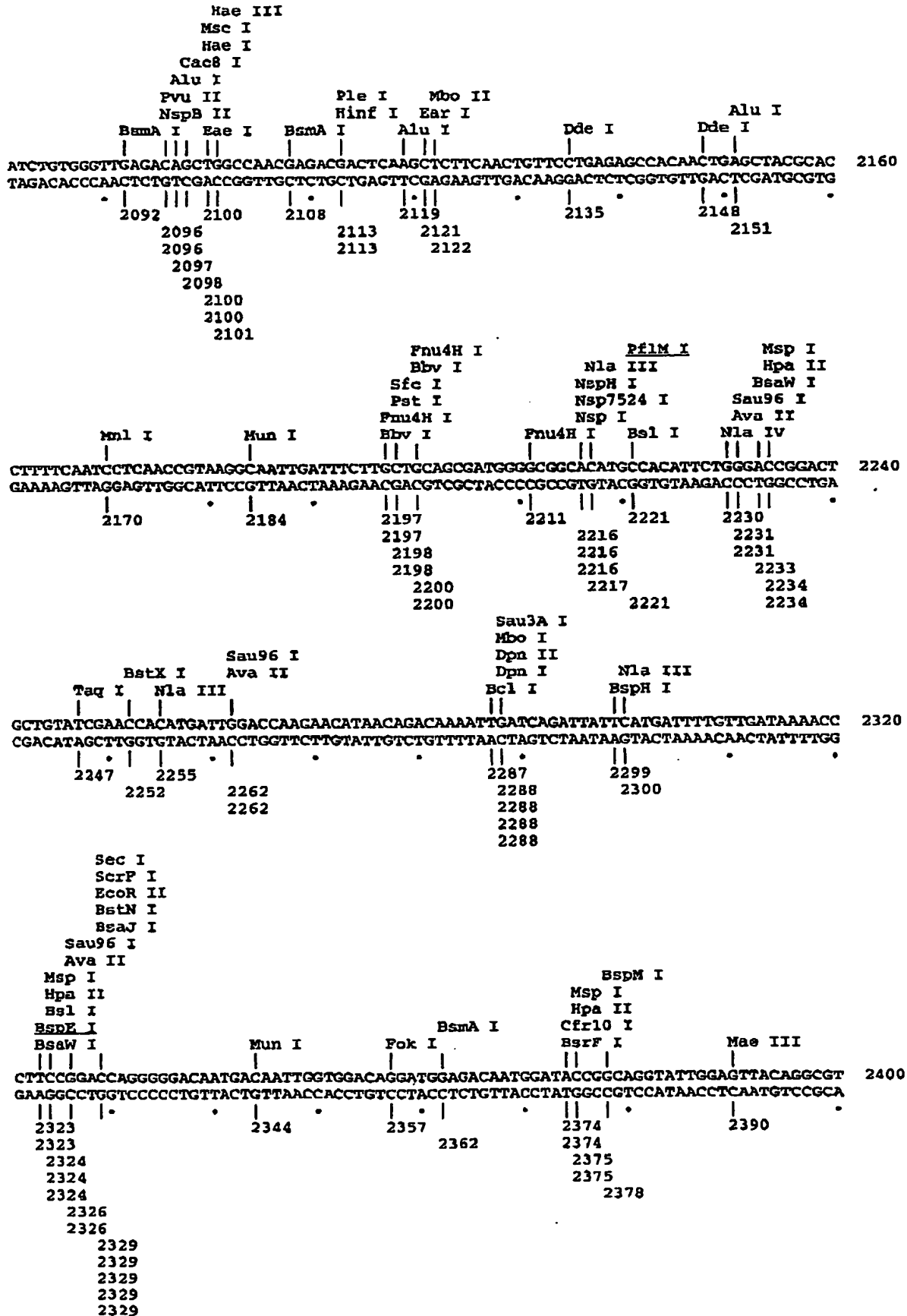


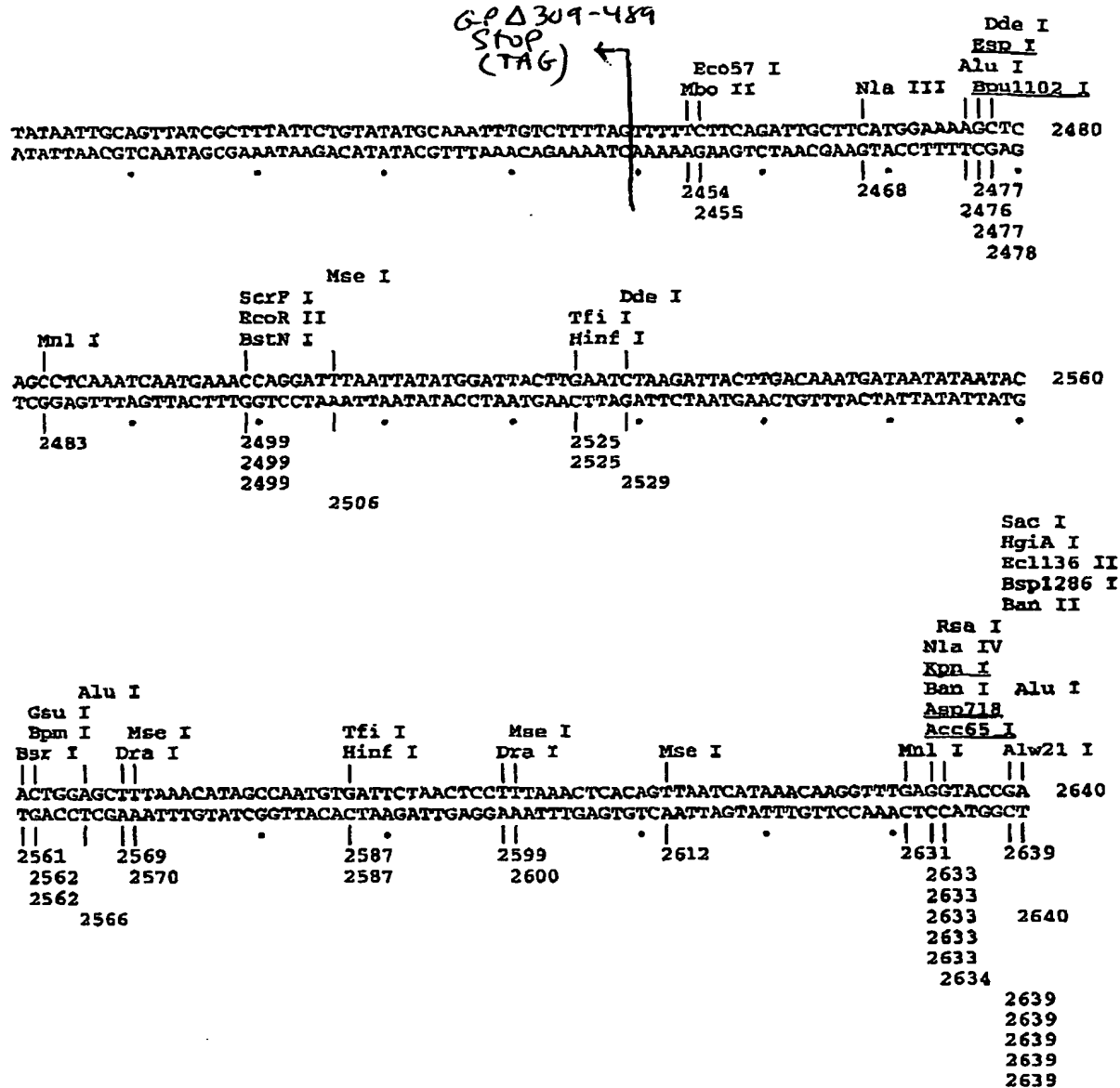
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Cla I  
Sau3A I  
Mbo I  
Dpn II  
BssS I Hinf I  
Mae III Ssp I Mnl I BspD I  
Mae III Dpn I Mnl I  
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962 970 979 983 989 990 993  
984 989 989 989 990 991 993  
Fok I  
Ple I Hinf I  
Xmn I Fok I Hinf I  
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1097 1106 1097 1106 1098 1106  
Mae I  
Sau3A I  
Mbo I  
Dpn II  
Dpn I  
Fok I Mun I Dpn I  
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1128 1139 1146 1146 1146 1146 1151 1159 1159 1164 1188  
Msp I  
Hpa II  
BsaW I  
Sau96 I  
Ava II  
Eco57 I  
HgiE II  
Bsl I  
Alu I  
Nph I  
AAC TAAAGATGGGGCTTCAGGTCCGGTGTCCCAACCAAGGTGGTCAATTATGAAGCTGGTGAATGGGCTGAAAACGTCT 1280  
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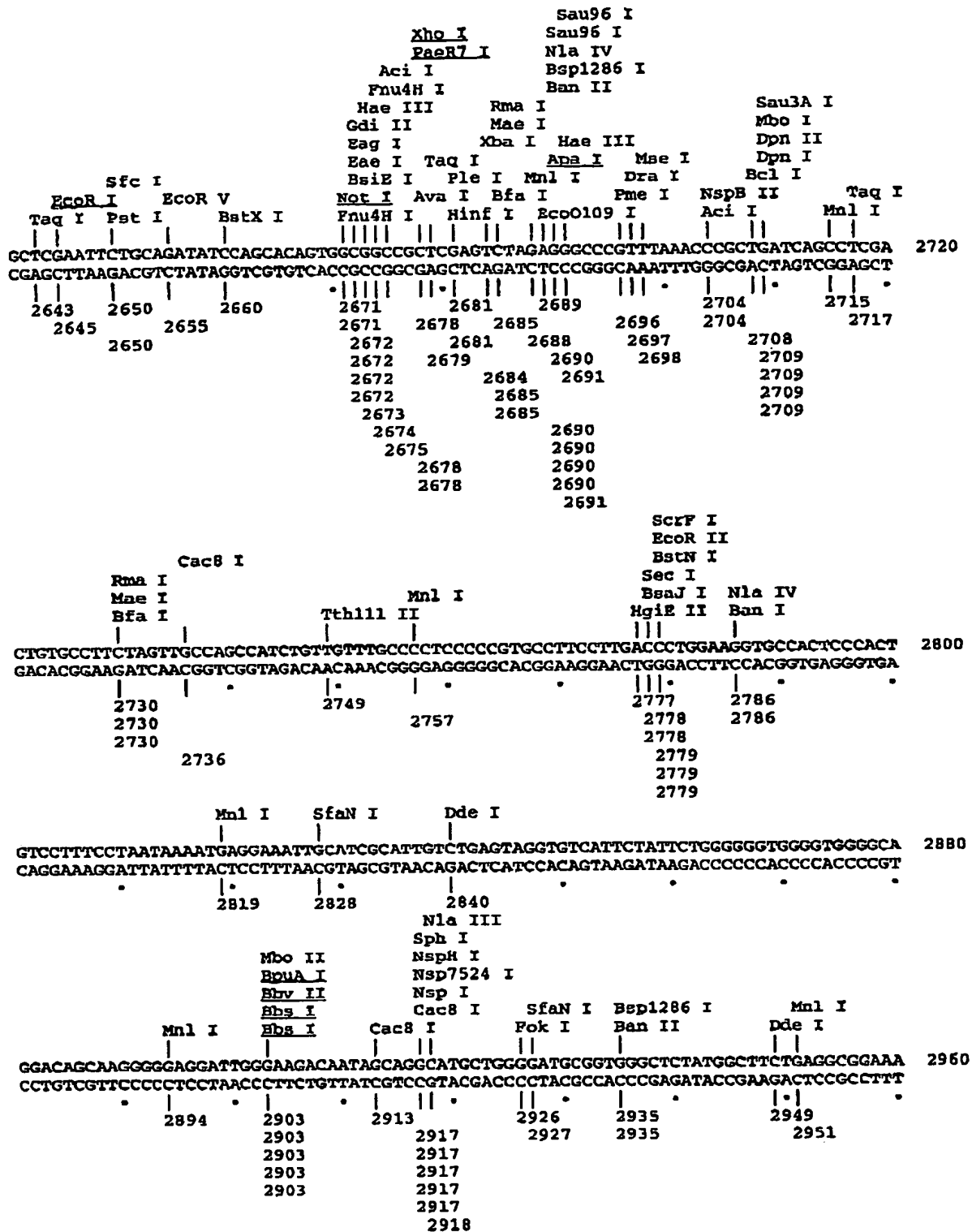












[illegible]



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3686 3694 3705 3729 3744  
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Scrf I  
Nci I  
Msp I  
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Bcn I  
Xma I  
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Ava I

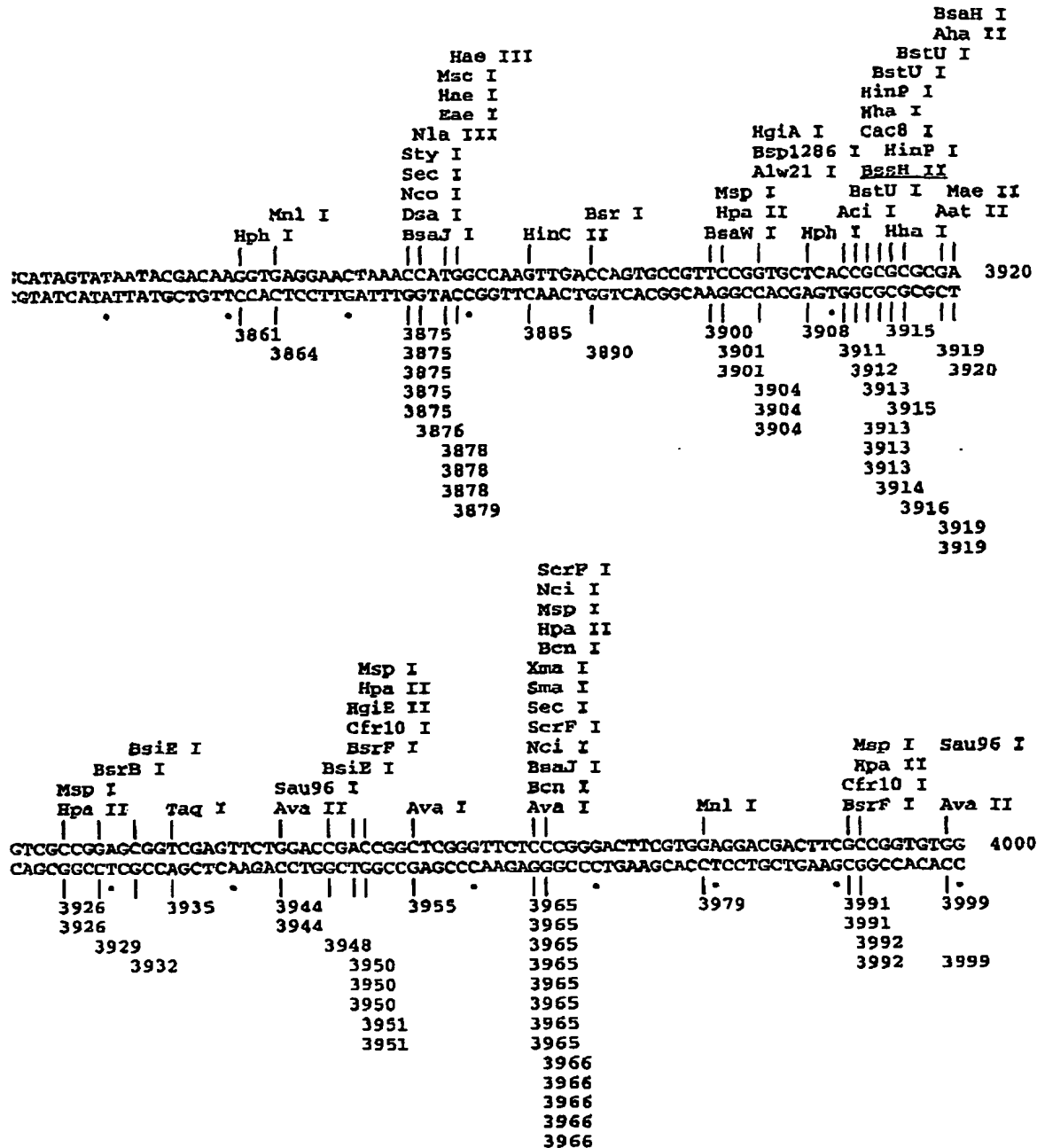
Alu I Alu I

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3765 3775 3794 3794 3795 3795 3795 3795 3795 3799 3800 3800 3800 3800 3806 3806 3807 3807 3811 3818 3819

Rna I  
Mae I  
Bfa I  
Sty I  
Sec I  
BsaJ I  
AVX II  
Hae III  
Stu I  
Hae I  
Mnl I  
BseR I  
Mnl I  
Mnl I  
Dde I  
Alu I  
Mnl I  
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Sec I  
Mnl I  
Fnu4H I  
Aci I

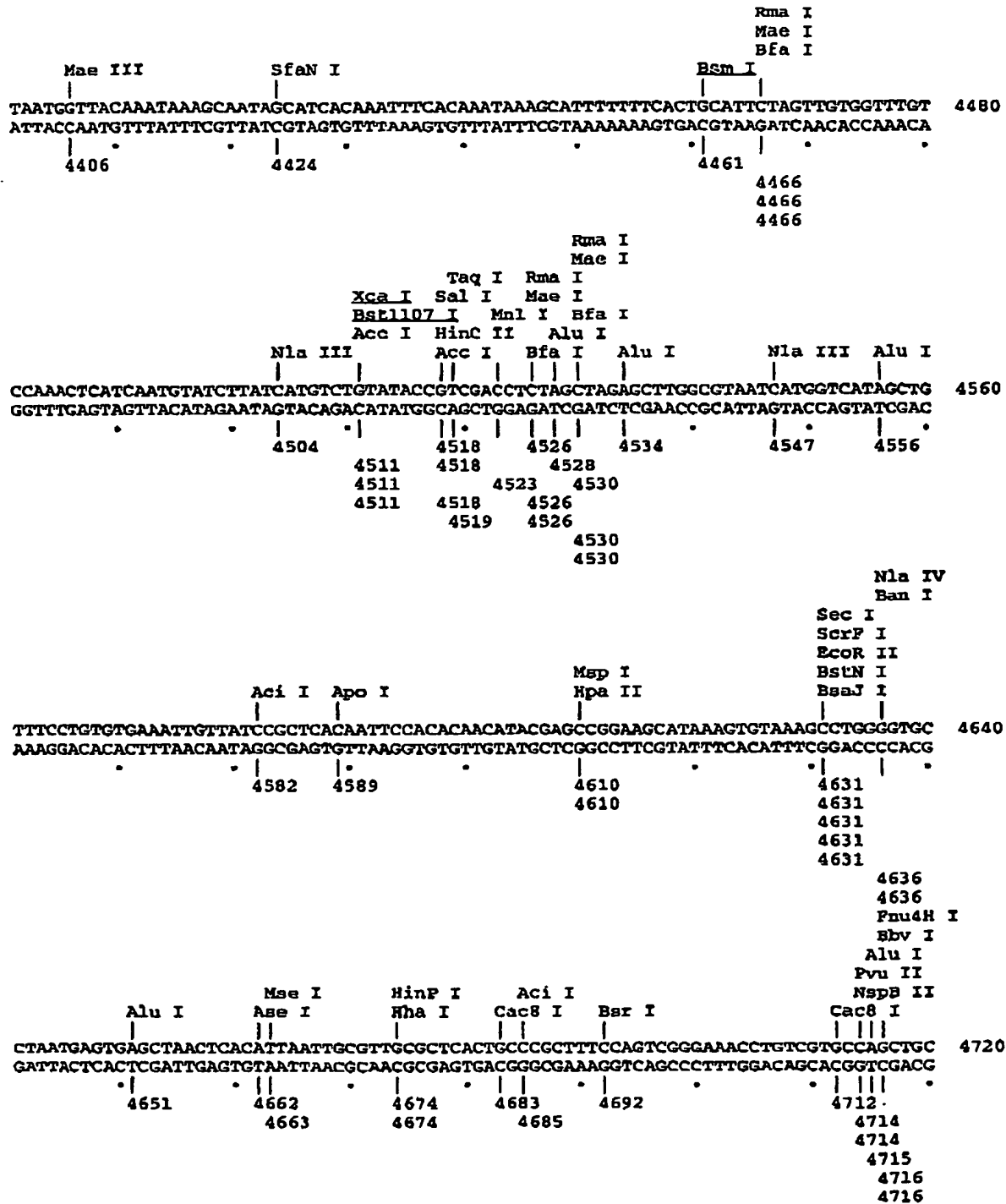
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Dpn I  
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Alw I  
Hinc II  
Mae II  
Afl III  
Pml I  
Eco72 I  
BsaA I  
Mse I  
Ase I



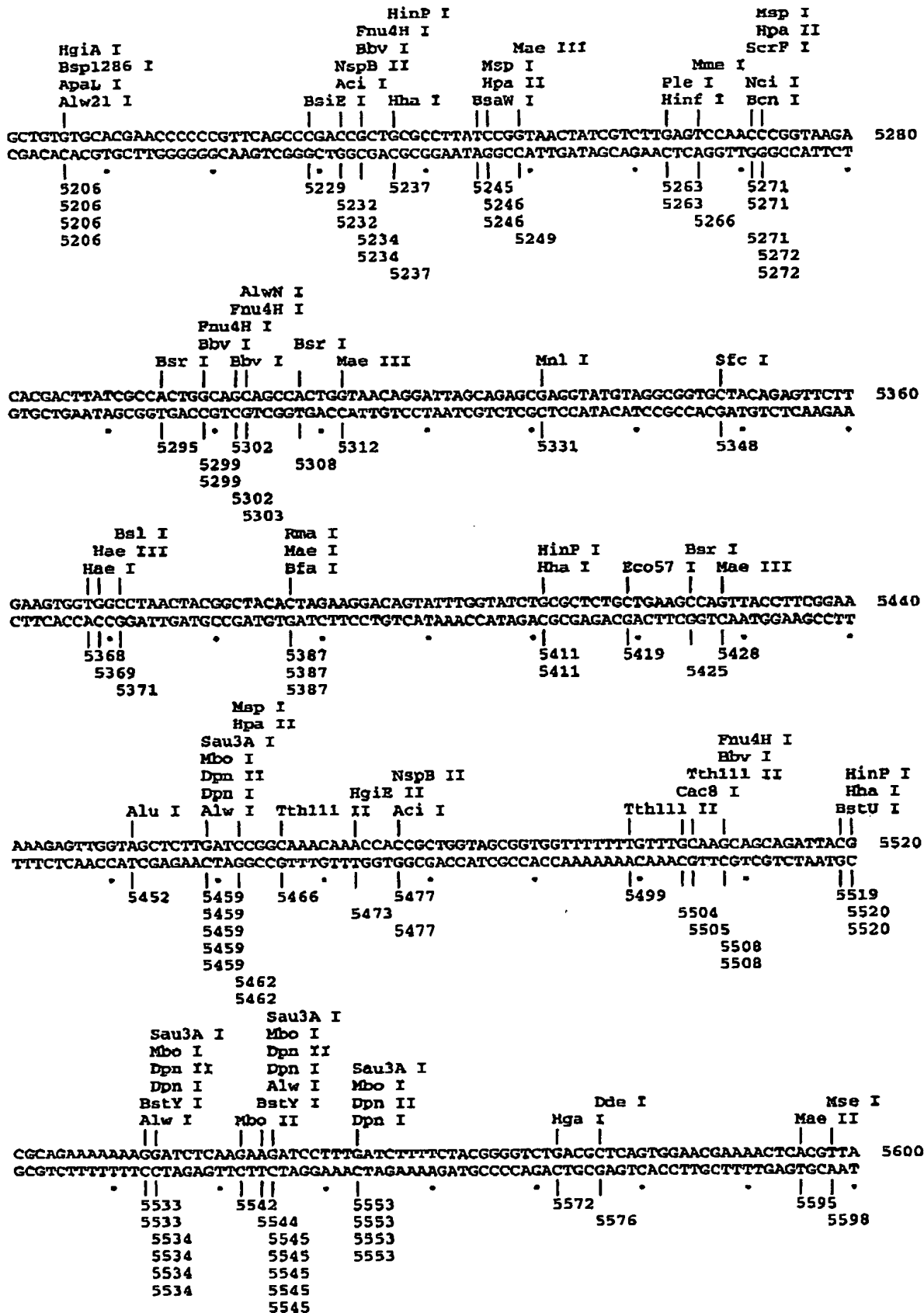


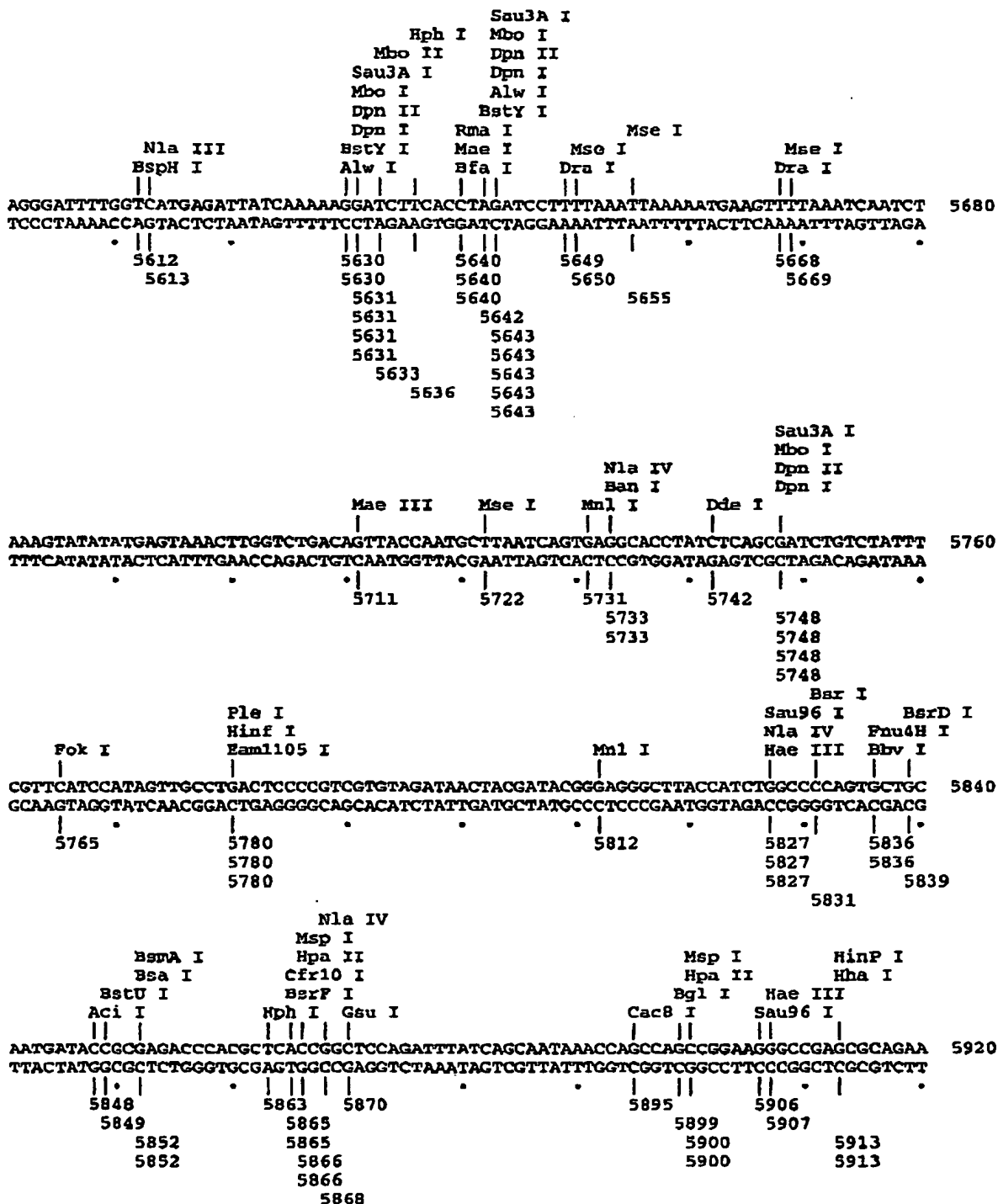


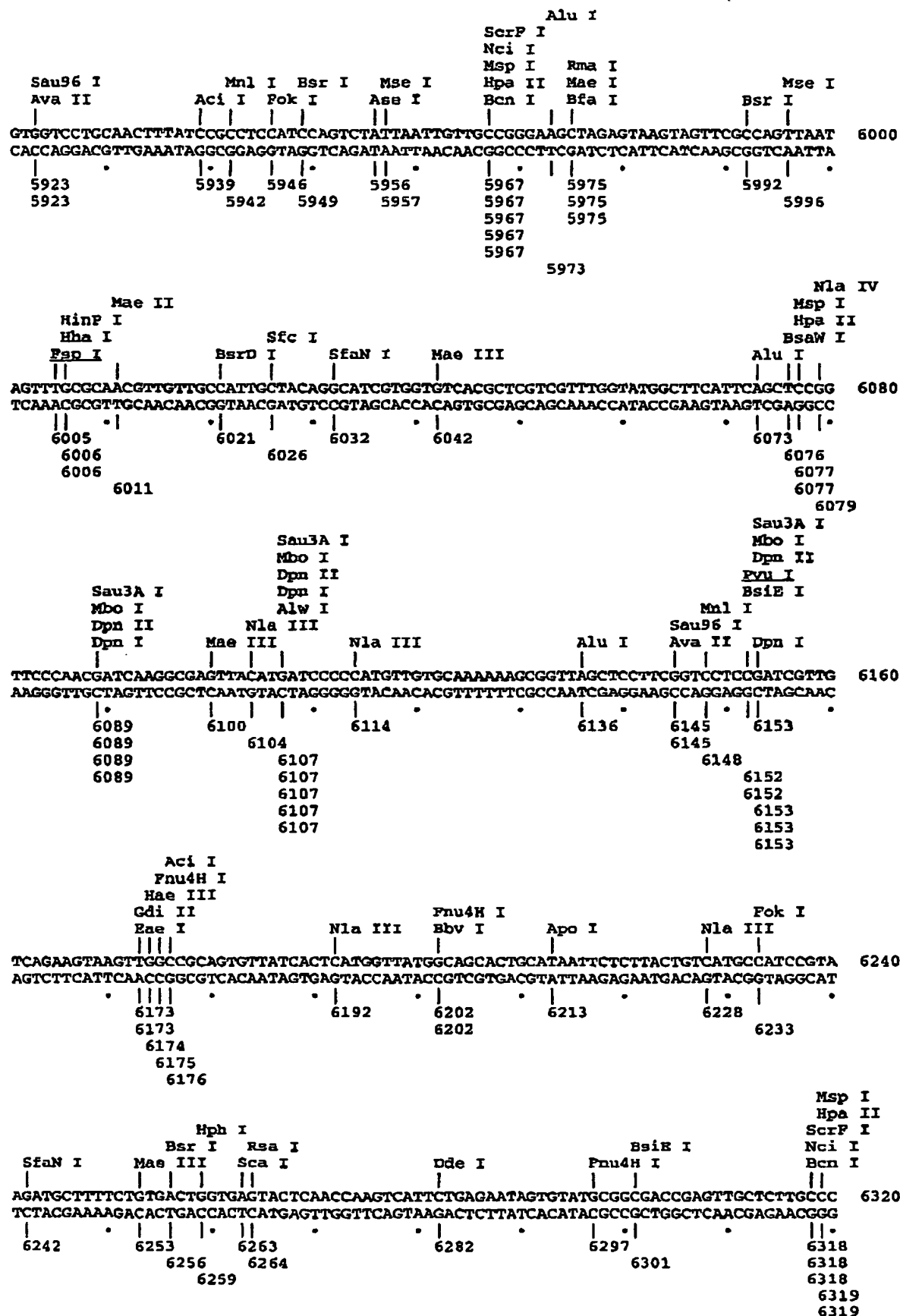
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NcoM I  
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Cfr10 I  
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BsrF I  
Gdi II  
Eag I  
Eae I  
Msp I  
ScrF I  
Nci I  
Bcn I  
BstU I Hae III  
Hinf I BsiE I  
Hha I Hpa II  
Dra III  
HgiA I  
Bsp1286 I  
ApaL I  
Alw21 I  
Cac8 I  
Mnl I  
BseR I  
Sec I  
BsaJ I  
Hae III  
Gdi II  
Eae I  
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4210  
4210  
4210  
4210  
4211  
4211  
Mae II  
Pml I  
Eco72 I  
BsaA I  
Afl III  
Tfi I  
Hinf I  
Taq I  
Aci I  
Aci I  
Pnu4H I  
Tfi I  
Hinf I  
Bcn I  
ScrF I  
Nci I  
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NcoM I  
Nae I  
Cfr10 I  
Cac8 I  
BsrF I  
Bsl I  
Gsu I  
Sau3A I  
Mbo I  
Dpn II  
BstY I  
Alw I  
Hga I  
BsaH I  
Aha II  
Fok I  
Mnl I  
BstU I  
Hha I  
Dpn I  
Nla III  
Mbo II  
Alu I  
Pnu4H I  
Bbv I  
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520







Aat II	6	Bsm I	1	Fnu4H I	33	Nsp7524 I	6
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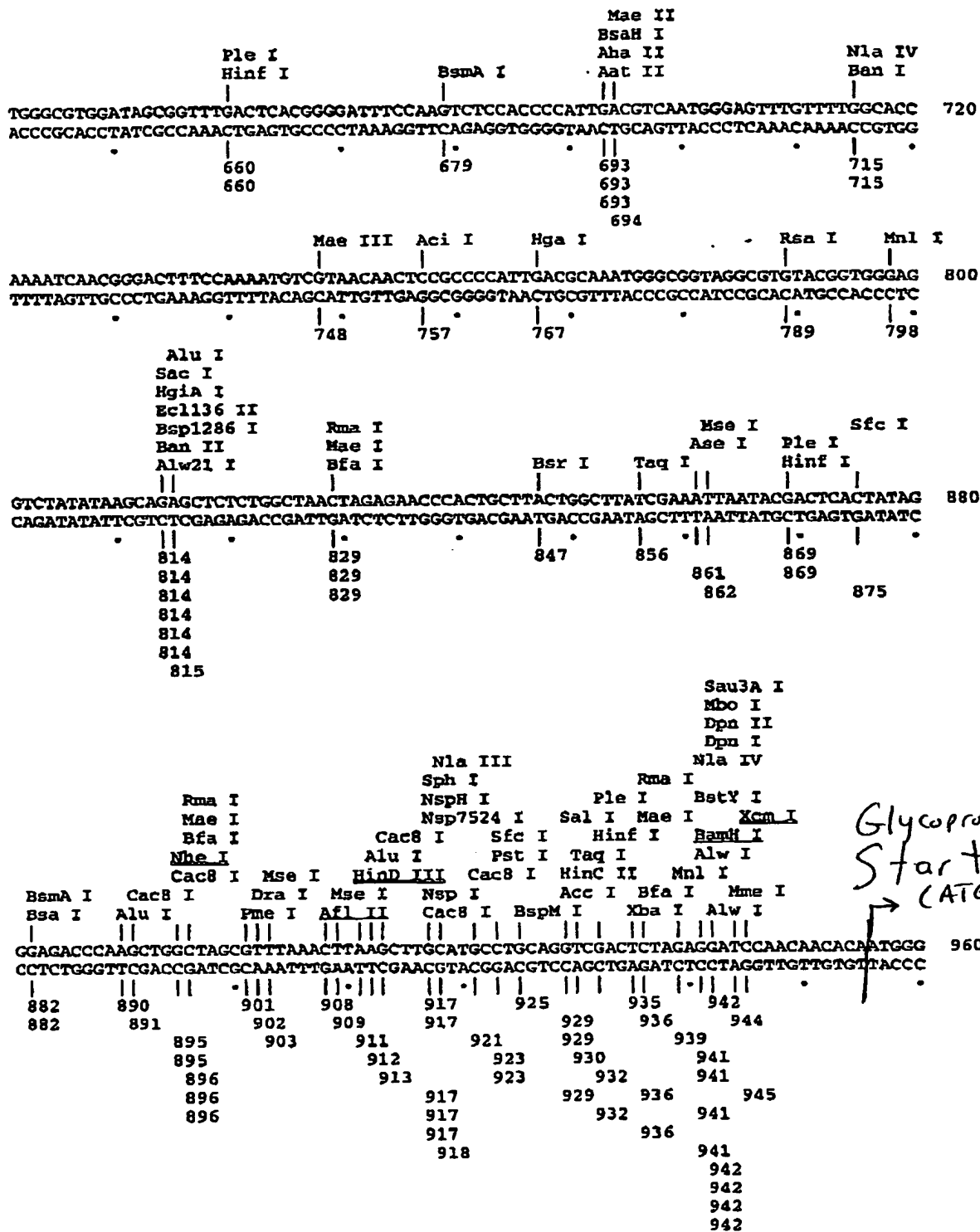
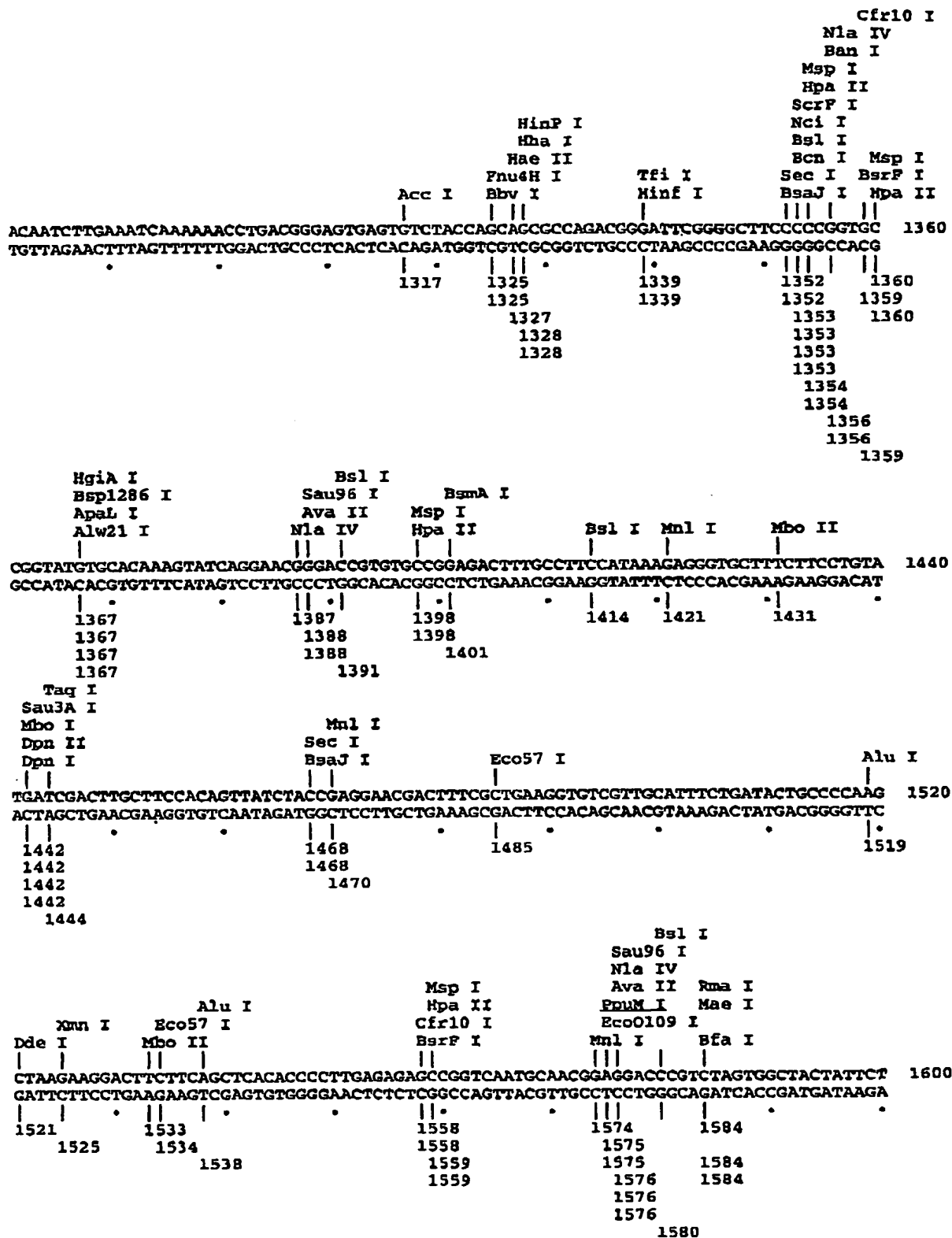
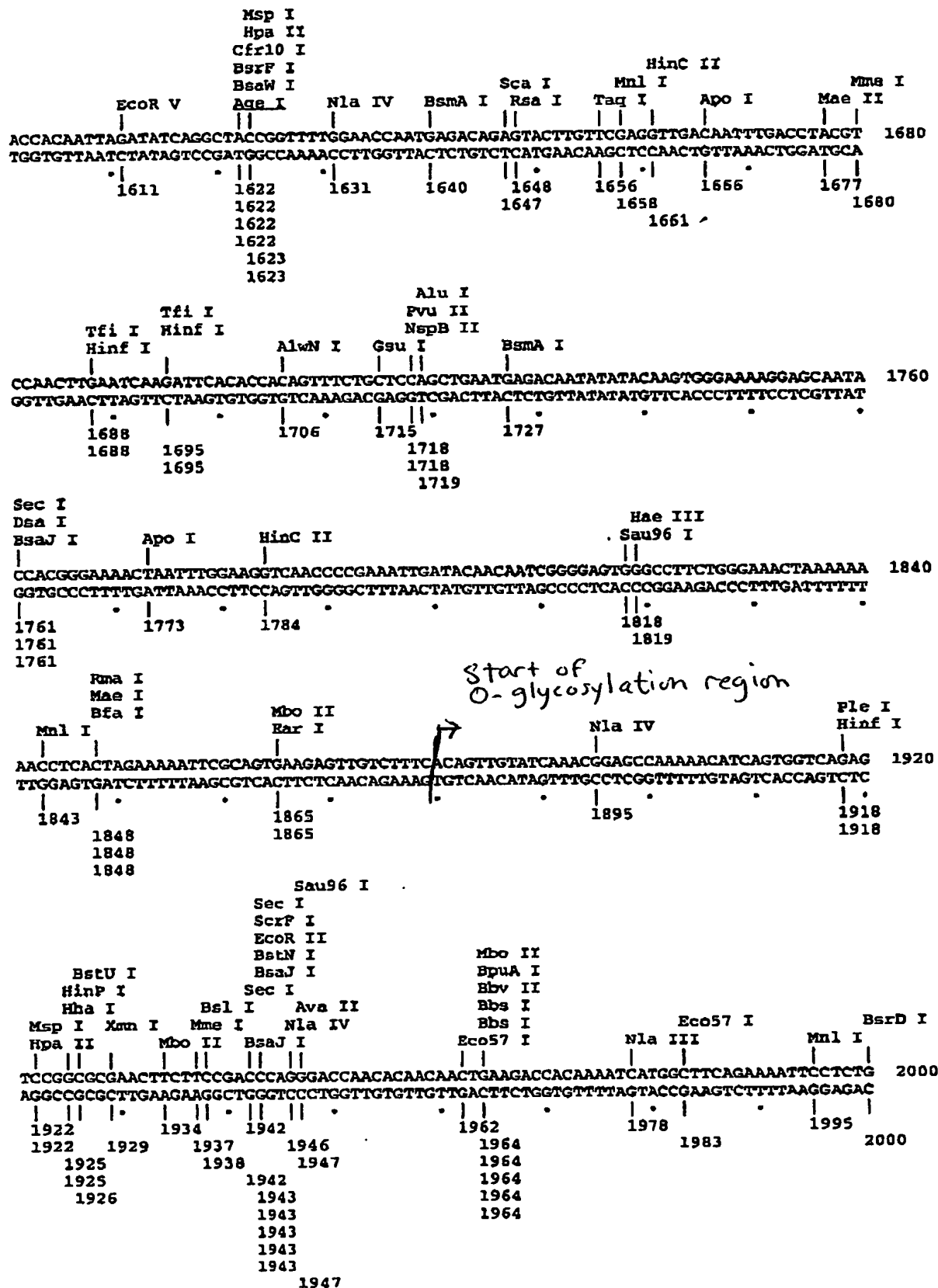


fig. 1C

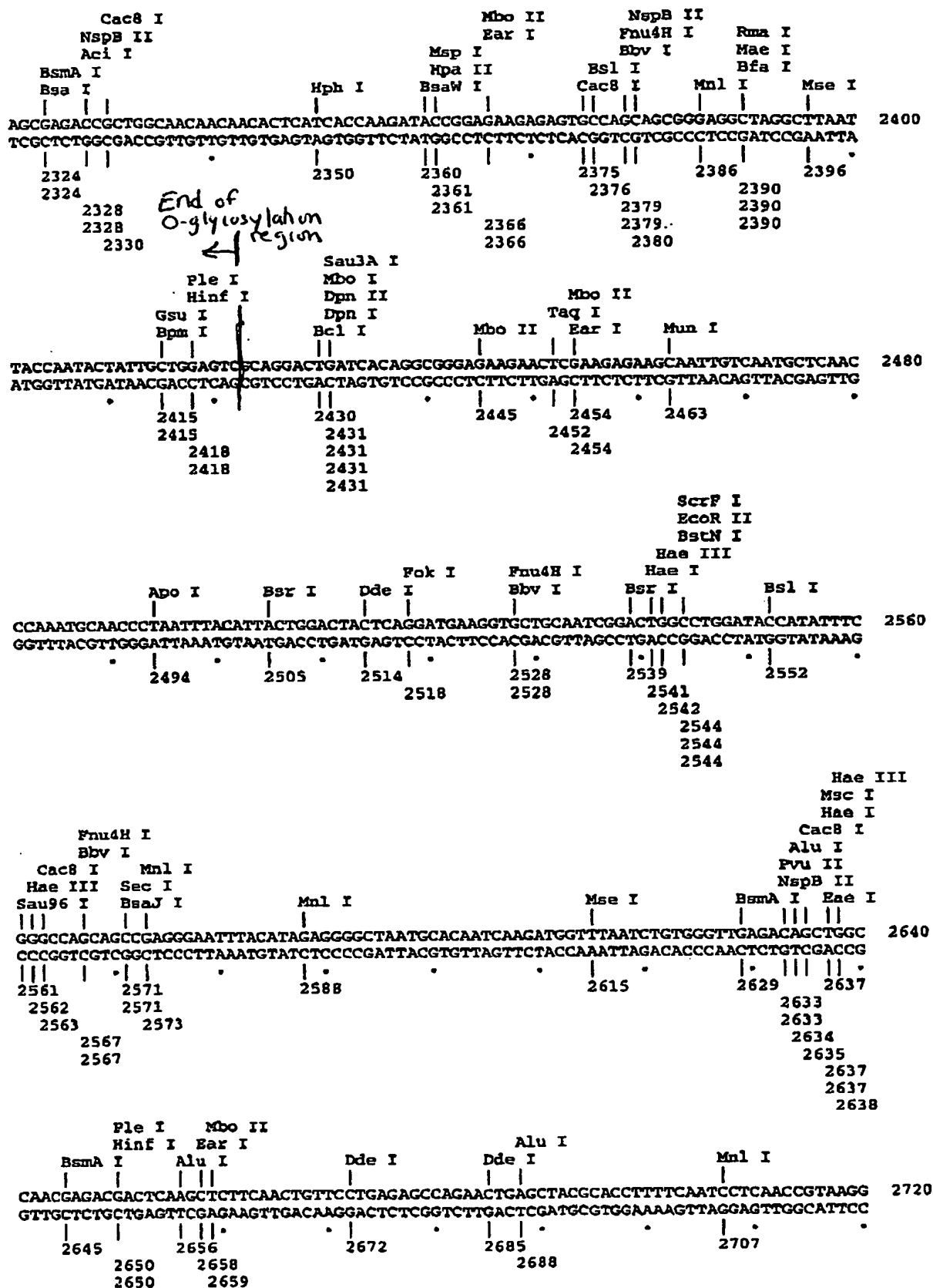


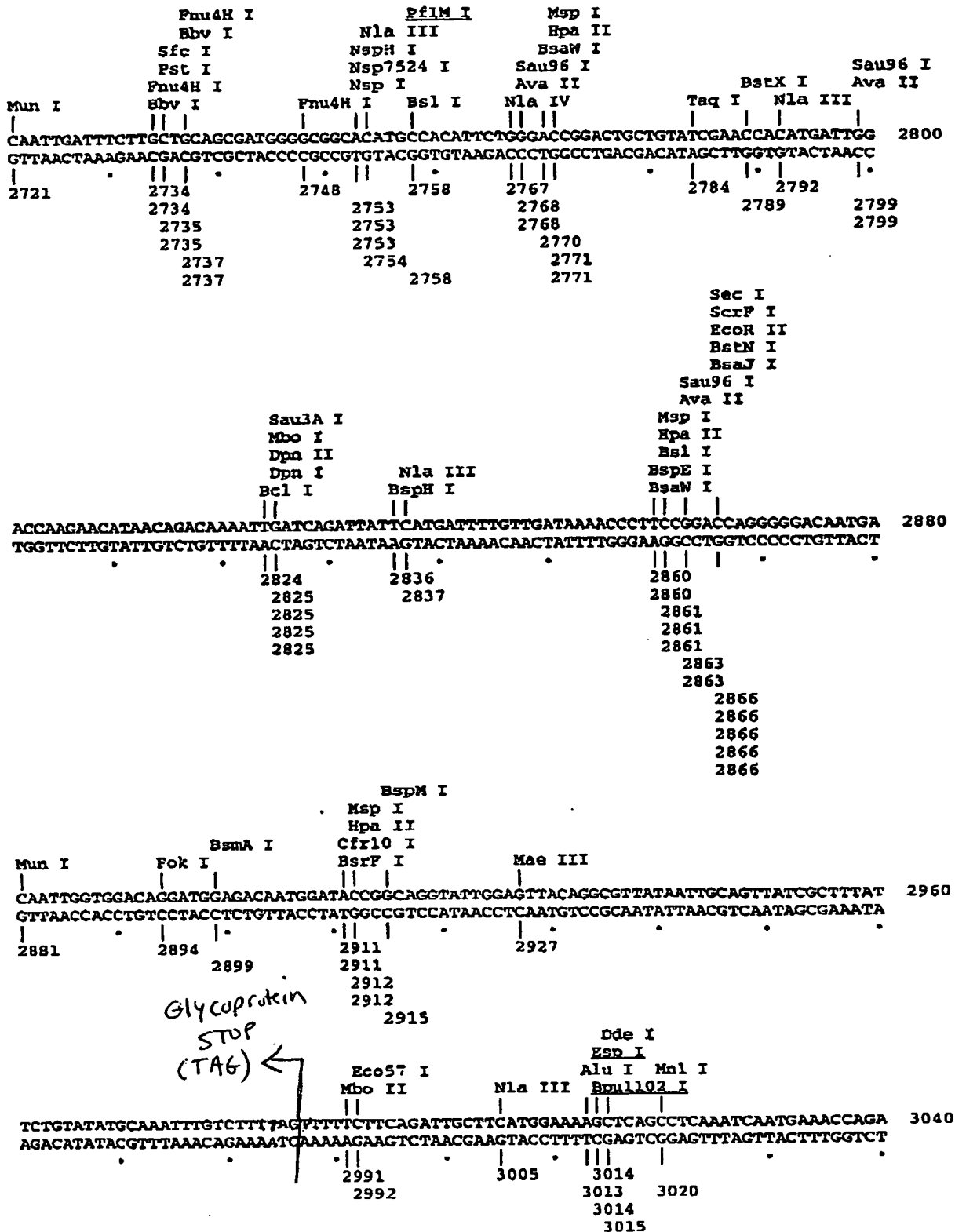
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962 970 979 983 989 1000  
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989  
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990  
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993  
Taq I Rna I  
Sal I Mae I  
Hinc II Bfa I  
Acc I Spe I Mae III  
Xmn I Pok I Ple I Hinf I  
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Sau3A I  
Mbo I  
Dpn II  
Taq I  
Fok I Mun I Dpn I Hinf I Mae II  
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BsaW I  
Sau96 I  
Ava II HgiE II  
Eco57 I Bsl I Hph I  
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 ApaI I  
 Alw21 I  
 Sfc I  
 Pst I  
 Fnu4H I  
 Bbv I  
 Alu I  
 SfaN I  
 Ple I  
 Hinf I  
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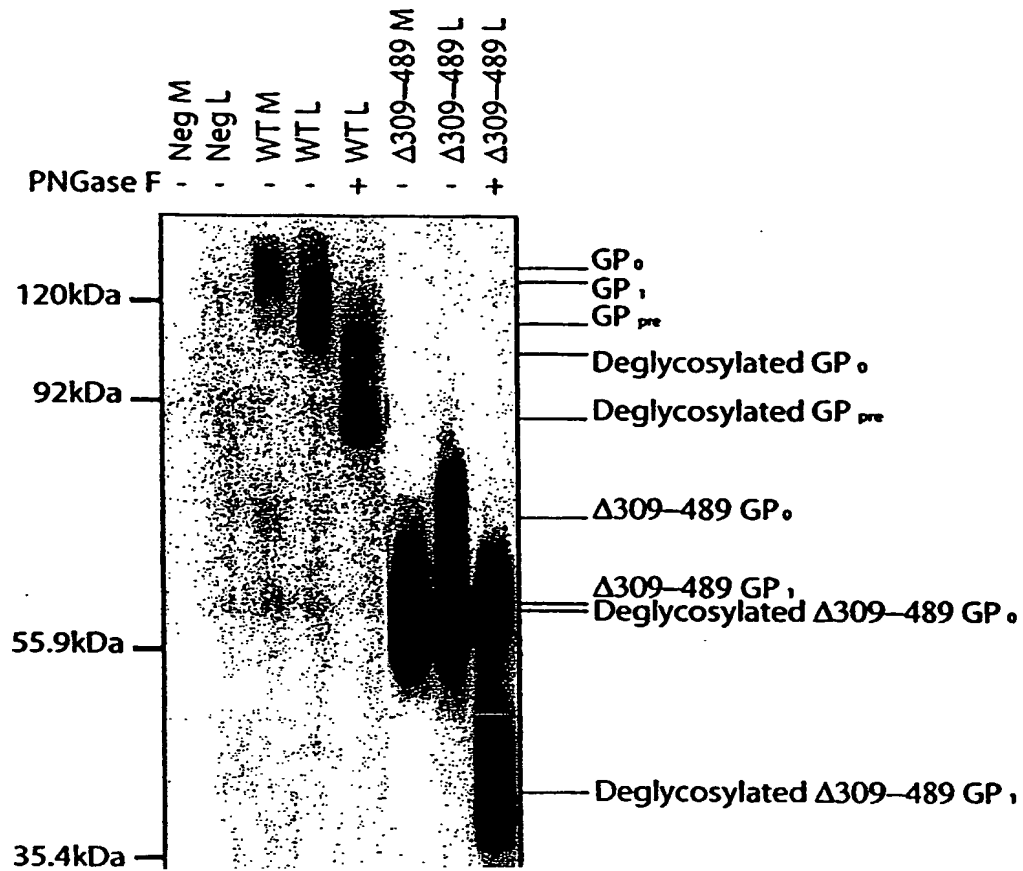
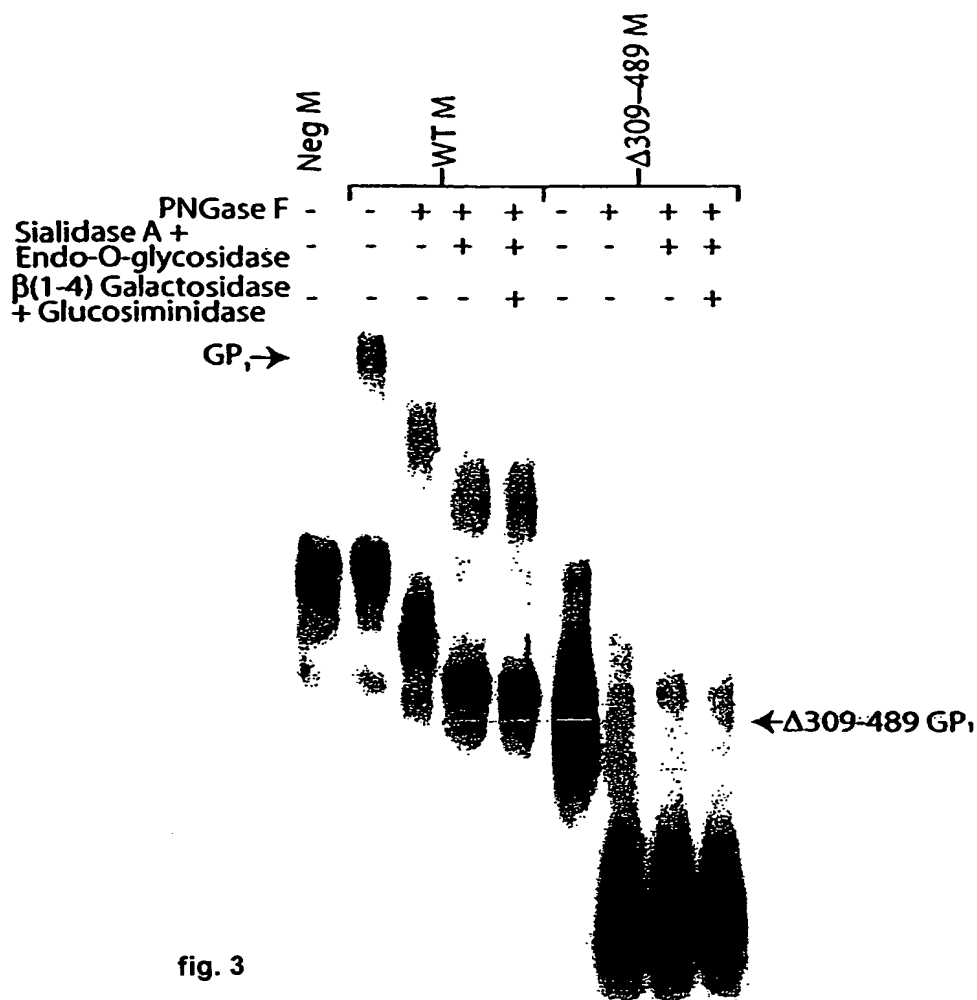


fig. 2





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